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☐ 1: AI732223. nf17g03.x5 NCI_CG...[gi:5053336]   ProbeSet, Taxonomy, LinkOut

```
LOCUS       AI732223                457 bp    mRNA    linear   EST 13-DEC-1999
DEFINITION  nf17g03.x5 NCI_CGAP_Pr1 Homo sapiens cDNA clone IMAGE:914068
            similar to contains element PTR7 repetitive element mRNA
            sequence.
ACCESSION   AI732223
VERSION     AI732223.1  GI:5053336
KEYWORDS    EST.
SOURCE      human.
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 457)
  AUTHORS   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
  TITLE     National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
            Tumor Gene Index
  JOURNAL   Unpublished (1997)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.
            , Michael Emmert-Buck, M.D., Ph.D.
            cDNA Library Preparation: David B. Krizman, Ph.D.
            cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
            DNA Sequencing by: Washington University Genome Sequencing Center
            Clone distribution: NCI-CGAP clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            www-bio.llnl.gov/bbrp/image/image.html

            This read is a RESEQUENCE of a previously sequenced human clone
            Original clone citation: National Cancer Institute, Cancer Genome
            Anatomy Project (CGAP), Tumor Gene Index
            This read has been verified (found to hit its original self in the
            correct orientation)
            Insert Length: 573    Std Error: 0.00
            Seq primer: -40UP from Gibco
            High quality sequence stop: 418.
FEATURES             Location/Qualifiers
     source          1..457
                     /organism="Homo sapiens"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:914068"
                     /clone_lib="NCI_CGAP_Pr1"
                     /sex="Male"
                     /dev_stage="45 years old"
                     /lab_host="DH10B"
                     /note="Vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st
                     strand cDNA was primed with oligo(dT)17 on 50 ng of
                     DNAse-treated, total cellular RNA obtained from 5,000-10
                     ,000 microdissected, histologically normal prostate
                     epithelial cells. Double-stranded cDNA was ligated to
                     EcoRI adaptors, 5 cycles of PCR applied to the cDNA with
                     an adaptor-specific primer, and the resulting PCR product
                     subcloned into pAMP10 by the UDG-cloning method (Life
                     Technologies). Average insert size is 600 bp. NOTE: Not
                     directionally cloned. This library was constructed by
                     David Krizman."
BASE COUNT       92 a      135 c      105 g      125 t
```